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Friday, Nov. 12<sup>th</sup> '54

Dear Rosalind,

Many thanks for your letter, and for the Fourier  $\eta$  certainly "looks better". I am just a little worried that the reversal of two terms of (as I remember)

opposite sign can change the origin so much. This is presumably due to the Lorentz factor bringing up the interstices as higher  $\lambda$ . Do you think that you are getting any diffraction effects? Perhaps an <sup>artificial</sup> temperature factor might help.

I agree with you about Casper's interpretation, but feel the matter will really remain open until you either get a

good picture of the protein lacking RNA, or with a  
good heavy atom (and even the latter may not make identification  
easy). If you are to cope with complex phases you may have  
to think in terms of two heavy atoms at different places  
(the SH and the tyrosine peptides) that would be very  
exciting.

I don't know anybody off-hand for your ARC job,  
but I'll ask Lohard. In our experience it usually  
takes a little time to find someone who is really suitable for  
this type of job.

Come again.

Yours.

Franklin